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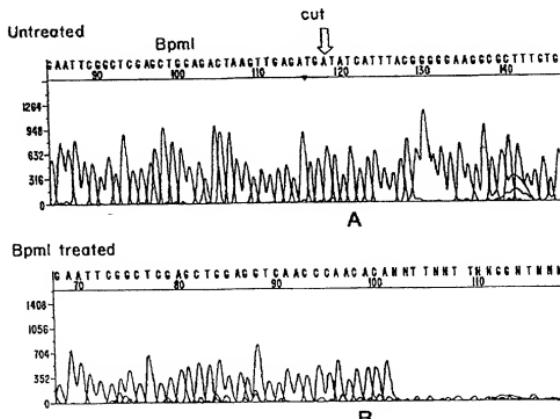
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(54) 350a: MULTIPLE SEQUENCING METHODS



(57) Abstract

The present invention provides a method for identifying a nucleic acid utilizing a run-off sequencing reaction of a relatively short portion of the nucleic acid. The method can be utilized, for example, to identify an EST from only a small portion of the EST and in an analysis of nucleotide polymorphisms. The figure depicts an untreated and a BpmI-treated sequencing reaction.